

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extra
035607	BMPR2_MOUSE	Mus musculus	Bone morphogenetic protein receptor type-2	35.498311	NaN	T379;S586;S680;S681;S843	34678516;22645316	MTSSLHRPFRVPWLLWAVLLVSTTA ASQNERLCAFKDPYQDGLIGESR ISHENGTILCSKSGSTCYGLWEKSG DINLVKQGCWSHIGDPQECHYECCV VTTTPPSIQNGTYRFCCSSTDLNCV NFTENFPPDPTTPLSPPHSFNRDETI HALASVSVLAVLIVALCFGYRMLTGD RKQGLHSMNMMEAAAAEPLDLD NLKLELIGRGRYGAVYKGS�DERPV AVKVFSFANRQNFINEKNIYRVPLM EHDNIARFIVGDERLTADGRMEYLL VMEYYPNGSLCKYLSLHTSDWVSSC RLAHSVTRGLAYLHTELPRGDHYKP AISHRDLNSRNLVKNDGACVISDF GLSMRLTGNRLVRPGEEDNAAISEV GTIRYMAPEVLEGAVNLRDCESALK QVDMYALGLIYWEVFMRCTDLFPGE SVPDYQMAFQTEVGNHPTFEDMQV LVSREKQRPKFPEAWKENS�AVRSL KETIEDCWDQDAEARLTAQCAEER MAELMMIWERNKSVSPVTPMSTA MQNERNLSHNRRVPKIGPYPDYSSS SYIEDSIHHTDSIVKNISSEHSMSST PLTIGEKNRNSINERQQAARIPSP ETSVTSLSTNTTTTNTTGLTPSTGMT TISEMPYPDETHLHATNVAQSIGPTP VCLQLTEEDLETNKLDPKVDKNLK ESSDENLMEHSLKQFSGPDLSSST SSLLYPLIKLAVEVTGQDFTQAANG QAQLIPDVPPAQIYPLPKQQLPKRP TSLPLNTKNSTKEPRLKFGNKHKSN LKQVETGVAKMNTINAAEPHVTVT MNGVAGRSHNVNSHAATTQYANGA VPAGQAANIVAHRSQEMLQNFQFIGE DTRLNINSSPDEHEPLLRREQQAGH DEGLDRLVDRRRRPLEGGRTNSN NNNSNPCSEQDILTQGVSTAADPG PSKPRRAQRPNLDSLATNILDGSSI QIGESTQDGKSGSGEKIKRRVKTPYS LKRWRPSTWVISTEPLDCEVNNNGS DRAVHKSSTAVYLAEGGTATTVSK DIGMNCL	False	True	2.173	3.64	2.305	1.884	1.339	5.0	3.149